

FIGURE 1

Figure 2B: Western blot showing APP processing in BACE1<sup>+/+</sup> and BACE1<sup>-/-</sup> cells. Lanes are labeled BACE1<sup>+/+</sup> and BACE1<sup>-/-</sup>. Molecular weight markers (kDa) are indicated on the left: 172, 111, 14.9, 6.1, 3.1. Bands are labeled: Full length APP,  $\beta$ -CTF,  $\alpha$ -CTF,  $\text{A}\beta$ , p3.

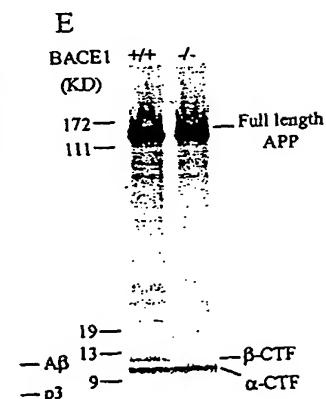
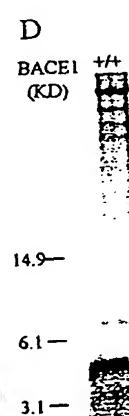
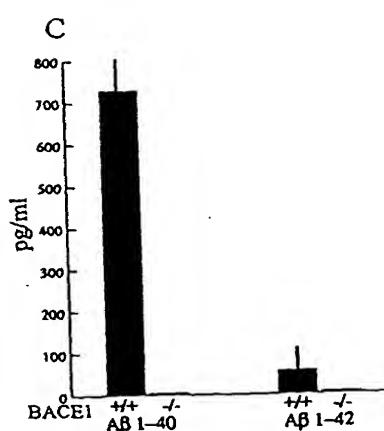
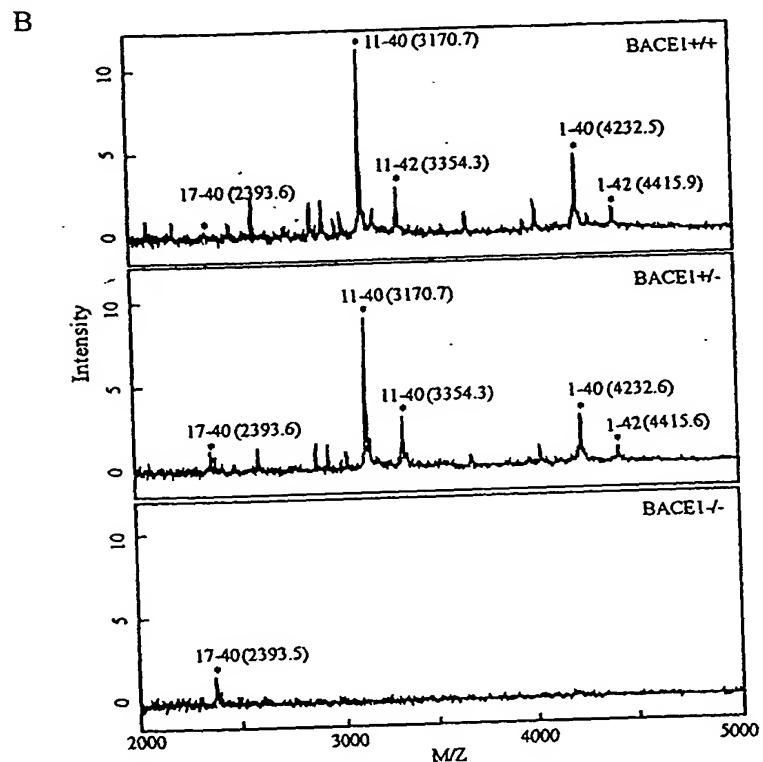
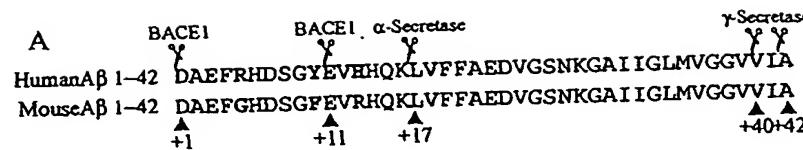


FIGURE 2

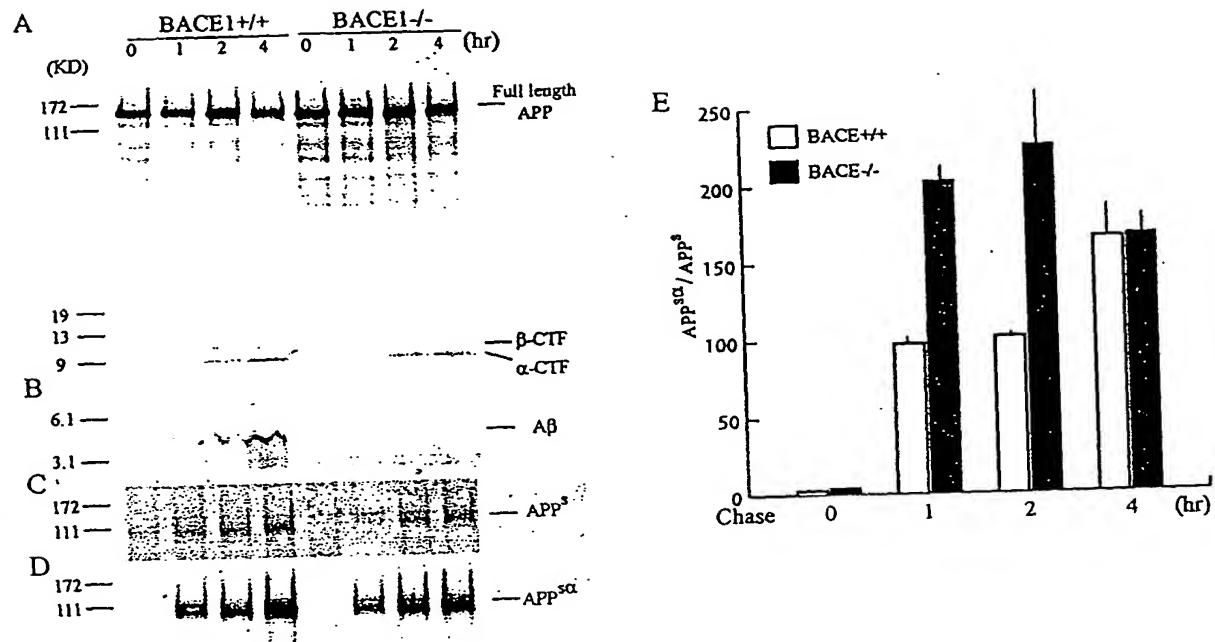


FIGURE 3

FIGURE 4 Dec 2000

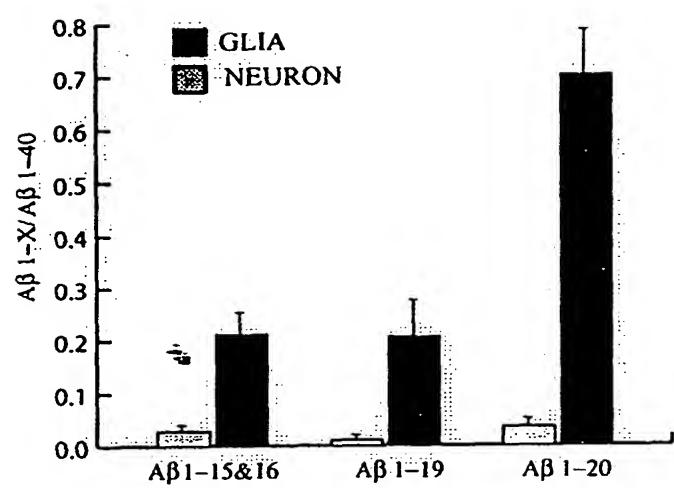


FIGURE 4